Fig.1

SEQ ID NO: 12

```
  1 GAATTCATGG TGTTTTGATC ATTTTAAATT TTTATATGGC GGGTGGTGGG CAACTCGCTT  60
 60 CCGGGCAACT CGCTTACCGA TTACGTTAGG GCTGATATTT ACGTAAAAAT CGTCAAGGGA 120
121 TGCAAGACCA AAGTAGTAAA ACCCCGGAGT CAACAGCATC CAAGCCCAAG TCCTTCACGG 180
181 AGAAACCCCA GCGTCCACAT CACGAGCGAA GGACCACCTC TAGGCATCGG ACGCACCATC 240
241 CAATTAGAAG CAGCAAAGCG AAACAGCCCA AGAAAAAGGT CGGCCCGTCG GCCTTTTCTG 300
301 CAACGCTGAT CACGGGCAGC GATCCAACCA ACACCCTCCA GAGTGACTAG GGGCGGAAAT 360
361 TTAAAGGGAT TAATTTCCAC TCAACCACAA ATCACAGTCG TCCCCGGTAT TGTCCTGCAG 420
421 AATGCAATTT AAACTCTTCT GCGAATCGCT TGGATTCCCC GCCCCTGGCC GTAGAGCTTA 480
481 AAGTATGTCC CTTGTCGATG CGATGTATCA CAACATATAA ATACTAGCAA GGGATGCCAT 540
541 GCTTGGAGGA TAGCAACCGA CAACATCACA TCAAGCTCTC CCTTCTCTGA ACAATAAACC 600
601 CCACAGAAGG CATTT                                                 615
```

Fig. 1 0

	SRE sequence		Amylase Activity (U/g dry mycelia)	(ratio)
SEQ ID NO: 6	**CGGAAATTTAAAGG**	taaP	1041	1
SEQ ID NO: 8	**CGGAAATTTAACGG**	taaS	1209	1.2
SEQ ID NO: 30	**CGGAAATTTAATTA**	MSRE2	792	0.8